

GenCore version 4.5
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OM protein - protein search, using sw model

Run on: March 1, 2001, 15:52:49 ; Search time 32.86 Seconds
(without alignments)
123.982 Million cell updates/sec

Title: US-09-331-631A-22_COPY_25_84

Perfect score: 350
Sequence: 1 EDDNNHHHGKSGQCVRRRC.....EKROERSRHEADRRSGEGSS 60

Scoring table: BLOSUM62
Gapop 10.0 , Gapext 0.5

Searched: 195891 seqs, 67900655 residues

Total number of hits satisfying chosen parameters: 195891

Minimum DB seq length: 0
Maximum DB seq length: 200000000

Post-processing: Minimum Match 0%
Maximum Match 100%

Listing first 45 summaries

Database : PIR_66:***
1: pirl:***
2: pirl2:***
3: pirl3:***
4: pirl4:***

Pred. No. is the number of results predicted by chance to have a score greater than or equal to the score of the result being printed, and is derived by analysis of the total score distribution.

SUMMARIES

Result No.	Score	Query Match	Query length	DB ID	Description
1	350	100.0	122	2 C53234	globulin-10 - maiz
2	350	100.0	582	2 B53234	vicilin-like stora
3	346	98.9	540	2 S21825	vicilin-like stora
4	346	98.9	573	2 A53234	globulin-15, GBLIS
5	342	97.7	407	2 T02258	globulin-1 - maize
6	303	86.6	236	2 T01662	globulin-1 - maize
7	129.5	37.0	637	2 S35221	globulin Bgl1 prec
8	96.5	27.6	1465	2 T23056	hypothetical prote
9	81.5	23.3	605	2 S06398	alpha-globulin typ
10	78	22.3	494	2 A48133	pre-mRNA splicing
11	74	20.1	971	2 T24866	hypothetical prote
12	71.5	20.4	997	2 T28872	hypothetical prote
13	71	20.3	699	2 A54660	histidine rich cal
14	70	20.0	144	2 F71446	hypothetical prote
15	70	20.0	2339	2 A42566	omega-conotoxin-se
16	69.5	19.9	241	2 S44893	PK1236.7 protein -
17	69.5	19.9	361	2 T39008	probable mating pr
18	69.5	19.9	549	2 T17525	proline-rich prote
19	69	19.7	407	1 EDBE03	immediate-early pr
20	69	19.7	588	1 FWCNAB	alpha-globulin B p
21	68.5	19.6	1407	1 S28589	trichohyalin B
22	68.5	19.6	2251	1 B54872	voltage-dependent
23	68.5	19.6	2270	2 A54972	voltage-dependent
24	67.5	19.3	466	2 T18575	cyclophilin isoform
25	67.5	19.3	1196	2 T14108	SH3-containing pro
26	67	19.1	385	2 T19201	hypothetical prote
27	67	19.1	509	2 S08059	alpha-globulin typ
28	67	19.1	566	2 S22477	vicilin precursor
29	67	19.1	897	2 T21688	hypothetical prote

30	66	18.9	2062	2 A59297	myosin X - mouse
31	66	18.9	2237	2 T45115	N-type calcium cha
32	65.5	18.7	480	2 T44225	hypothetical prote
33	65.5	18.7	796	2 JC7355	peroxisome prolif
34	65.5	18.7	2222	2 A37490	voltage-dependent
35	65.5	18.7	2272	2 C54972	voltage-dependent
36	65	18.6	57	2 T16083	hypothetical prote
37	65	18.6	107	2 A29995	protamine p2 precu
38	65	18.6	1105	2 T47582	hypothetical prote
39	65	18.6	1130	2 T30251	hypothetical prote
40	65	18.6	1276	2 T02711	repetin - mouse
41	64	18.3	241	2 T16849	hypothetical prote
42	64	18.3	640	2 A41726	hypothetical prote
43	63.5	18.1	423	2 T43929	homeotic protein B
44	63.5	18.1	520	2 S08237	DnaJ protein homol
45	63.5	18.1	763	2 T27620	legumin A2 precurs
					hypothetical prote

ALIGNMENTS

RESULT 1
C53234
globulin-10 - maize
C:Species: Zea mays (maize)
C:Date: 02-May-1994 #sequence_revision 18-Nov-1994 #text_change 29-Oct-1999
C:Accession: C53234; S21823
R:Belanger, F.C.; Kriz, A.L.
Genetics 129, 863-872, 1991
A:Title: Molecular basis for allelic polymorphism of the maize Globulin-1 gene.
A:Reference number: A53234; MUID:92090707
A:Contents: Black Beauty popcorn
A:Accession: C53234
A:Status: preliminary
A:Molecule type: DNA
A:Residues: 1-122 <BFL>
A:Cross-references: EMBL:X59085; NID:q22281; PIDN:CAA41811.1; PID:q22282
A:Note: sequence extracted from NCBI backbone (NCBIN:71282, NCBIPI:71286)

Query Match 100.0%; Score 350; DB 2; Length 122;
Best Local Similarity 100.0%; Pred. No. 6.5e-29;
Matches 60; Conservative 0; Mismatches 0; Indels 0; Gaps 0;

OY 1 EDDNNHHHGKSGQCVRRRCEDRPWHORPCTECREEREKROERSRHEADRRSGEGSS 60
Db 25 EDDNNHHHGKSGQCVRRRCEDRPWHORPCTECREEREKROERSRHEADRRSGEGSS 84
RESULT 2
B53234
vicilin-like storage protein Gbl1-L, embryo - maize
N:Alternate names: globulin-1L
C:Species: Zea mays (maize)
C:Date: 02-May-1994 #sequence_revision 18-Nov-1994 #text_change 24-Nov-1999
C:Accession: B53234; S21824
R:Belanger, F.C.; Kriz, A.L.
Genetics 129, 863-872, 1991
A:Title: Molecular basis for allelic polymorphism of the maize Globulin-1 gene.
A:Reference number: A53234; MUID:92090707
A:Accession: B53234
A:Status: preliminary
A:Molecule type: DNA
A:Residues: 1582 <BFL>
A:Cross-references: EMBL:X59083; NID:q22283; PIDN:CAA41809.1; PID:q22284
A:Experimental source: Inbred line W64A6
A:Note: sequence extracted from NCBI backbone (NCBIPI:71285)
C:Genetics:
A:Gene: Gbl1-L
A:Introns: 167/1; 225/3; 252/3; 349/3
C:Superfamily: glycinin

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Query Match          100.0%; Score 350; DB 2; Length 582;
Best Local Similarity 100.0%; Pred. No. 2.5e-28;
Matches 60; Conservative 0; Mismatches 0; Indels 0; Gaps 0;

QY 1 EDDNNHHHGKSGQCVRCEDRPMHQRPRCLDQCREEEREKROERSRHEADDRSGGSS 60
|||||
Db 25 EDDNNHHHGKSGQCVRCEDRPMHQRPRCLDQCREEEREKROERSRHEADDRSGGSS 84

RESULT 3
S21825
vicillin-like storage protein G1b1-S, embryo - maize
C:Species: Zea mays (maize)
C:Date: 20-Feb-1995 #sequence_revision 20-Feb-1995 #text_change 11-Jan-2000
C:Accession: S21825
R:Kriz, A.L.
Submitted to the EMBL Data Library, April 1991
A:Reference number: S21823
A:Accession: S21825
A:Status: preliminary
A:Molecule type: DNA
A:Residues: 1-540 <R1>
A:Cross-references: EMBL:X59084; NID:g22285; PIDN:CAA41810.1; PID:g22287
C:Genetics:
A:Gene: G1b1-S
A:introns: 170/1; 195/2; 222/2; 319/2
C:Superfamily: glycIn

Query Match          98.9%; Score 346; DB 2; Length 540;
Best Local Similarity 98.3%; Pred. No. 6e-28;
Matches 59; Conservative 1; Mismatches 0; Indels 0; Gaps 0;

QY 1 EDDNNHHHGKSGQCVRCEDRPMHQRPRCLDQCREEEREKROERSRHEADDRSGGSS 60
|||||
Db 25 EDDNNHHHGKSGQCVRCEDRPMHQRPRCLDQCREEEREKROERSRHEADDRSGGSS 84

RESULT 4
A53234
globulin-1S, GLB1S - maize
C:Species: Zea mays (maize)
C:Date: 02-May-1994 #sequence_revision 18-Nov-1994 #text_change 11-Jan-2000
C:Accession: A53234; A43642
R:Belanger, F.C.; Kriz, A.L.
Genetics 129, 863-872, 1991
A:Title: Molecular basis for allelic polymorphism of the maize globulin-1 gene.
A:Reference number: A53234; MUID:92090707
A:Accession: A53234
A:Status: preliminary
A:Molecule type: DNA
A:Residues: 1-573 <BEL>
A:Experimental source: Inbred line Va 26
A:Note: sequence extracted from NCBI backbone (NCBIN:71280, NCBI:P:71284)
R:Kriz, A.L.
Biochem. Genet. 27, 239-251, 1989
A:Title: Characterization of embryo globulins encoded by the maize Glb genes.
A:Reference number: A43642; MUID:89374022
A:Accession: A43642
A:Status: preliminary
A:Molecule type: protein
A:Residues: 87-100 <KR1>
C:Superfamily: glycIn

Query Match          98.9%; Score 346; DB 2; Length 573;
Best Local Similarity 98.3%; Pred. No. 6.3e-28;
Matches 59; Conservative 1; Mismatches 0; Indels 0; Gaps 0;

QY 1 EDDNNHHHGKSGQCVRCEDRPMHQRPRCLDQCREEEREKROERSRHEADDRSGGSS 60
|||||
Db 25 EDDNNHHHGKSGQCVRCEDRPMHQRPRCLDQCREEEREKROERSRHEADDRSGGSS 84
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RESULT 5
T02258
globulin1 - maize
C:Species: Zea mays (maize)
C:Date: 05-Mar-1999 #sequence_revision 05-Mar-1999 #text_change 29-Oct-1999
C:Accession: T02258
R:Bhatramakl, D.; Kriz, A.L.
Submitted to the EMBL Data Library, May 1995
A:Description: Nucleotide sequence analysis of a novel globulin1 null allele from the
A:Reference number: Z14643
A:Accession: T02258
A:Status: preliminary; translated from GB/EMBL/DBJ
A:Molecule type: DNA
A:Residues: 1-407 <BHA>
A:Cross-references: EMBL:U28017; NID:g927238; PIDN:AAB60295.1; PID:g927239
A:Experimental source: strain IHP
C:Genetics:
A:Gene: G1b1
A:introns: 168/1; 226/3; 254/1; 351/1

Query Match          97.7%; Score 342; DB 2; Length 407;
Best Local Similarity 96.7%; Pred. No. 1.2e-27;
Matches 58; Conservative 2; Mismatches 0; Indels 0; Gaps 0;

QY 1 EDDNNHHHGKSGQCVRCEDRPMHQRPRCLDQCREEEREKROERSRHEADDRSGGSS 60
|||||
Db 25 EDDNNHHHGKSGQCVRCEDRPMHQRPRCLDQCREEEREKROERSRHEADDRSGGSS 84

RESULT 6
T01662
globulin-1 - maize (fragment)
C:Species: Zea mays (maize)
C:Date: 19-Feb-1999 #sequence_revision 19-Feb-1999 #text_change 11-Jan-2000
C:Accession: T01662
R:Hilton, H.; Gaut, B.S.
Genetics 150, 863-872, 1998
A:Title: Speciation and domestication in maize and its wild relatives: evidence from
A:Reference number: Z14386; MUID:98429537
A:Accession: T01662
A:Status: preliminary; translated from GB/EMBL/DBJ
A:Molecule type: DNA
A:Residues: 1-236 <HIL>
A:Cross-references: EMBL:AF064222; NID:g3414836; PIDN:AMC31465.1; PID:g3414837
A:Experimental source: subspecies parviflumis
C:Genetics:
A:introns: 166/1; 224/3
C:Superfamily: glycIn

Query Match          86.6%; Score 303; DB 2; Length 236;
Best Local Similarity 88.3%; Pred. No. 6.8e-24;
Matches 53; Conservative 1; Mismatches 0; Indels 6; Gaps 1;

QY 1 EDDNNHHHGKSGQCVRCEDRPMHQRPRCLDQCREEEREKROERSRHEADDRSGGSS 60
|||||
Db 25 EDDNNHHHGKSGQCVRCEDRPMHQRPRCLDQCREE-----EKSrHEADDRSGGSS 78

RESULT 7
S35221
globulin Beg1 precursor - barley
C:Species: Hordeum vulgare (barley)
C:Date: 03-Feb-1994 #sequence_revision 03-Feb-1994 #text_change 21-Jul-2000
C:Accession: S35221
R:Heck, G.R.; Chamberlain, A.K.; Ho, T.H.D.
Mol. Gen. Genet. 239, 209-218, 1993
A:Title: Barley embryo globulin 1 gene, Beg1: characterization of cDNA, chromosome ma
A:Reference number: S35221; MUID:93287988
A:Accession: S35221
A:Molecule type: mRNA
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